



# DESIGN AND FORMULATION ASPECTS OF MONOCLONAL ANTIBODIES FOR SARS-COV-2 TREATMENT

**KATARINA SMILKOV, PhD**

**Associate professor of Biopharmacy and Biotechnology**

**Division of Pharmacy/Faculty of Medical Sciences**

**University Goce Delcev Štip, N. Macedonia**

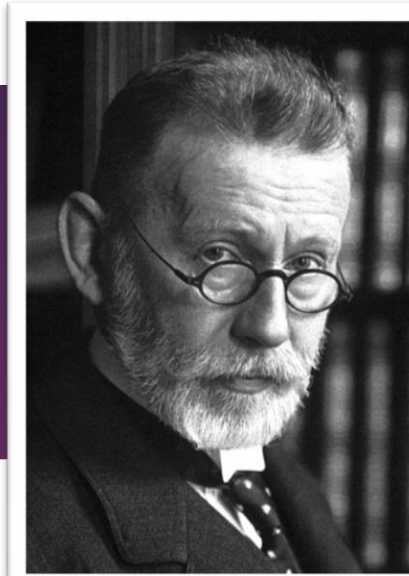
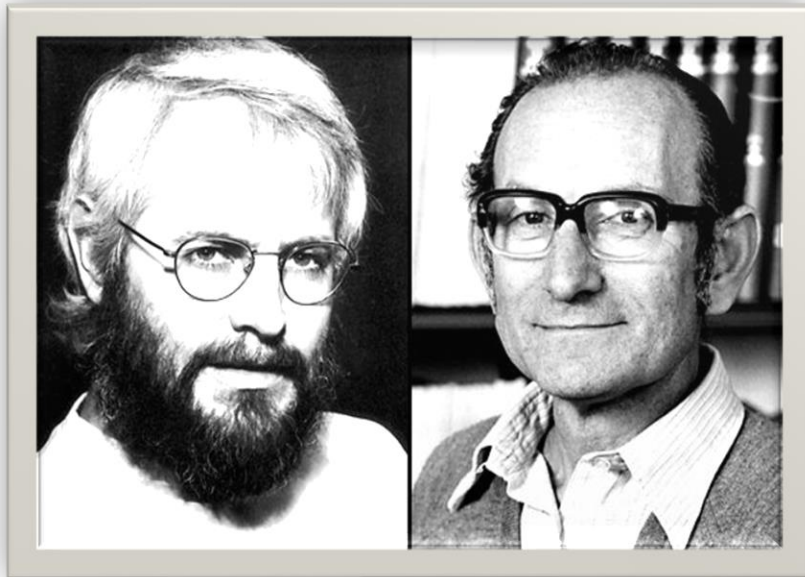
**[katarina.smilkov@ugd.edu.mk](mailto:katarina.smilkov@ugd.edu.mk)**



## Paul Erlich 1854 - 1915

Nobel Prize for Physiology or Medicine in 1908

- ▶ Side chain theory of immunity
- ▶ Antibodies - magic bullets 1899/1906

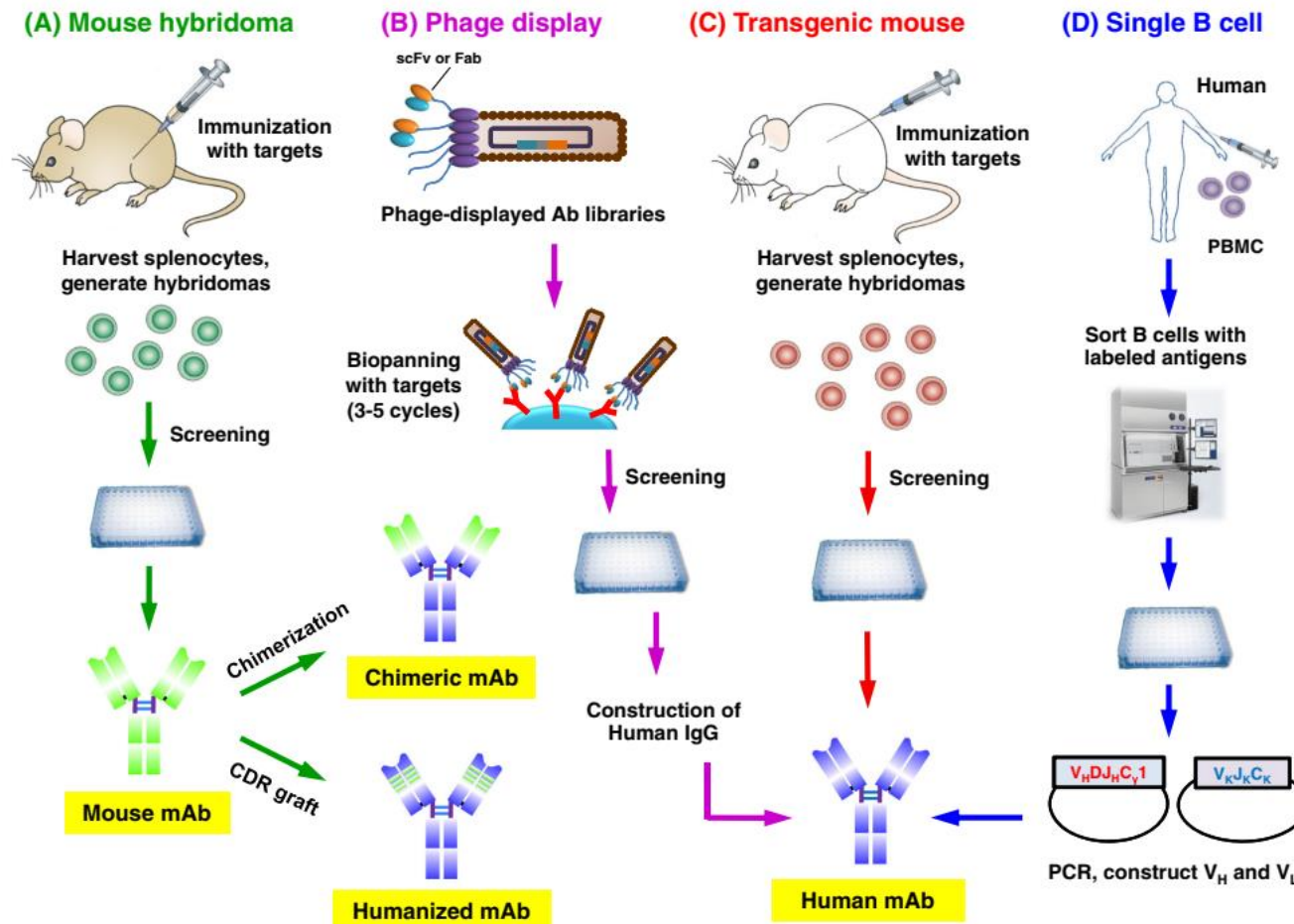


## Georges J.F Köhler and Cèsar Milstein

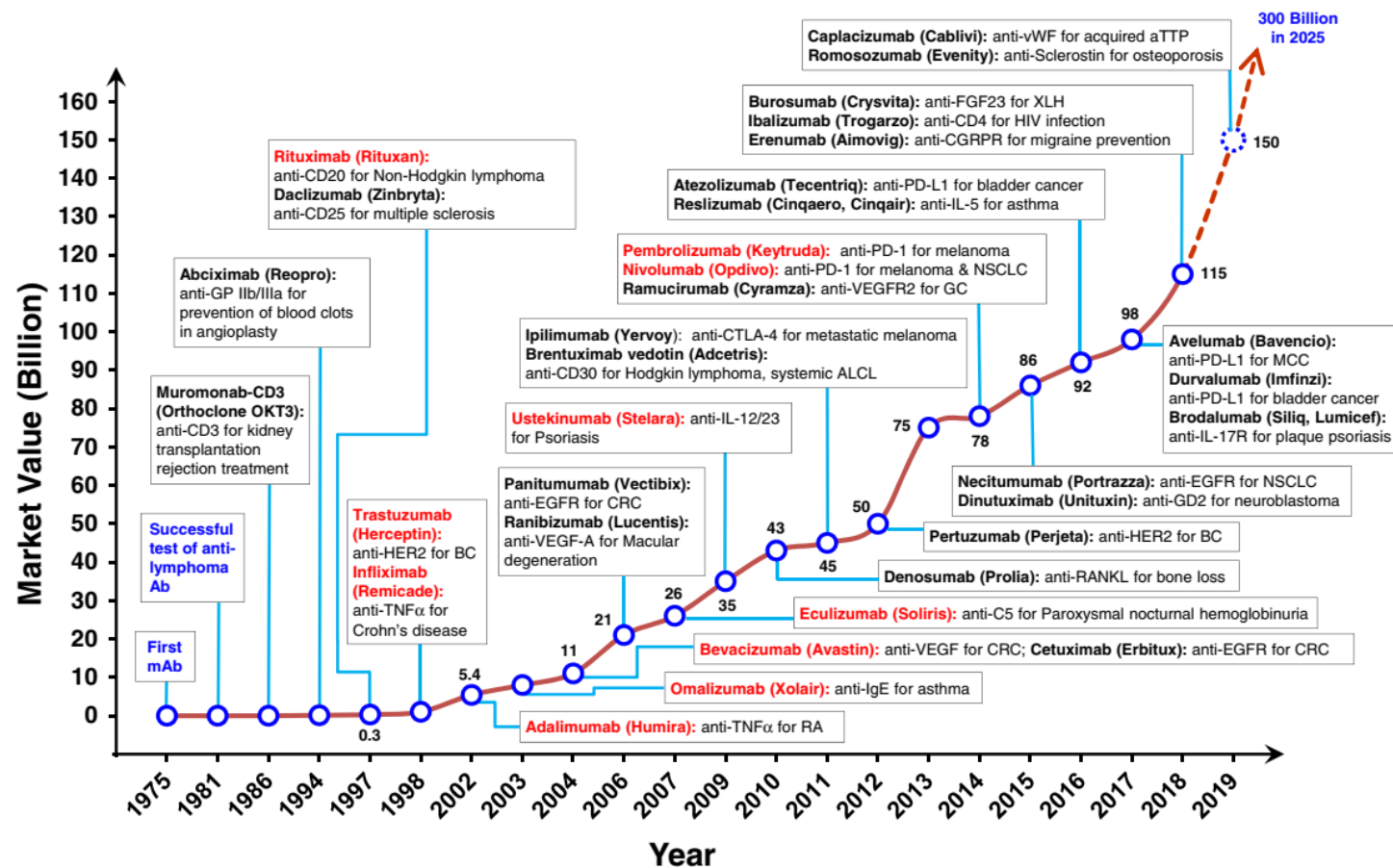
Nobel Prize for Physiology or Medicine in 1984

- ▶ 1975 introduced hybridoma technology
- ▶ the first monoclonal to be approved (OKT-3®, muromonab-CD3) was approved in 1986 for a single treatment regimen to protect from life-threatening tissue rejection following kidney transplant

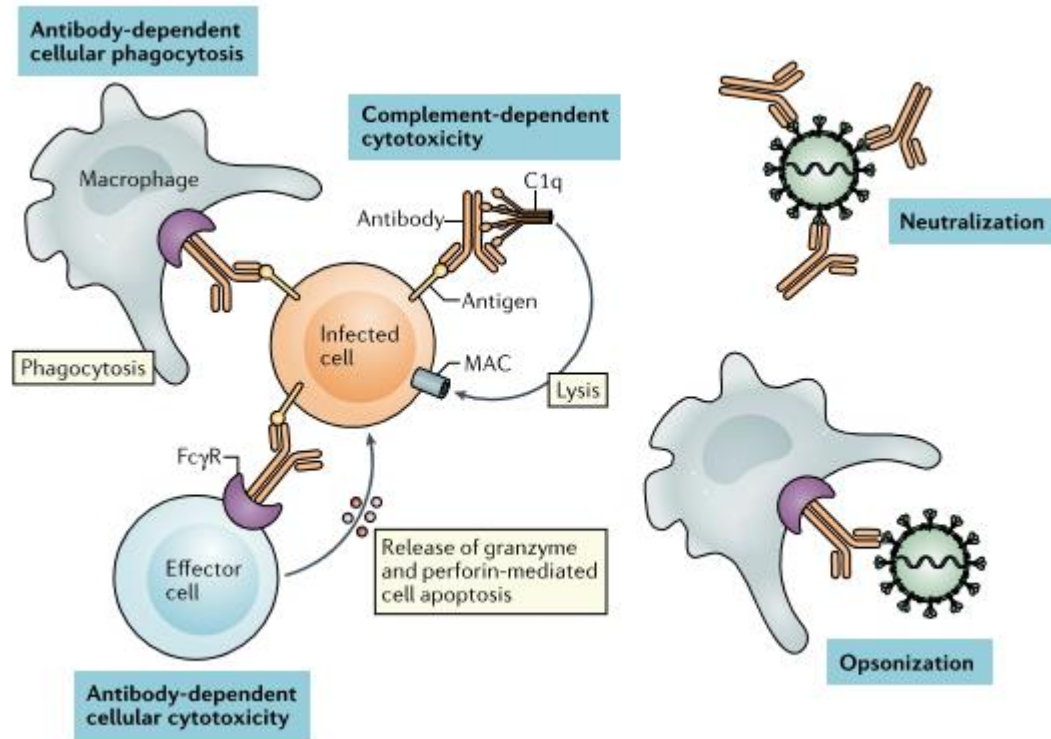
# Design of monoclonal antibodies (mAbs)



# mAbs industry is constantly growing



# mAbs mechanism of action



## Directly interfere with (viral) pathogenesis

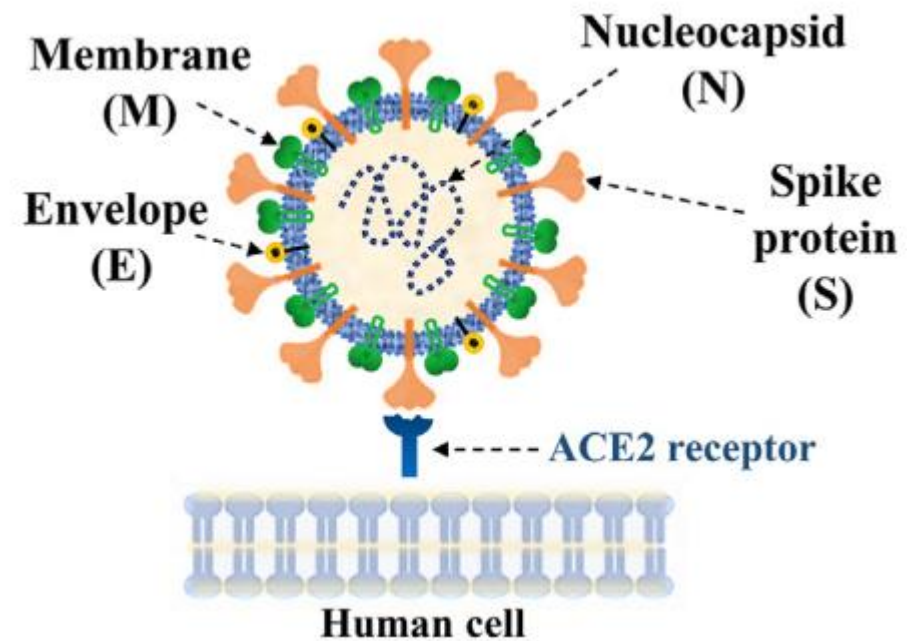
- ▶ binding of a neutralizing antibody to the virion can prevent target cell binding and/or fusion
  - ▶ antibody binding opsonizes the virions or infected cells for phagocytic uptake
  - ▶ mAbs can facilitate target cell death via
    - complement fixation and membrane attack complex (MAC) activation
    - antibody-dependent cytotoxicity
- resulting in apoptosis or necrosis of the infected cell



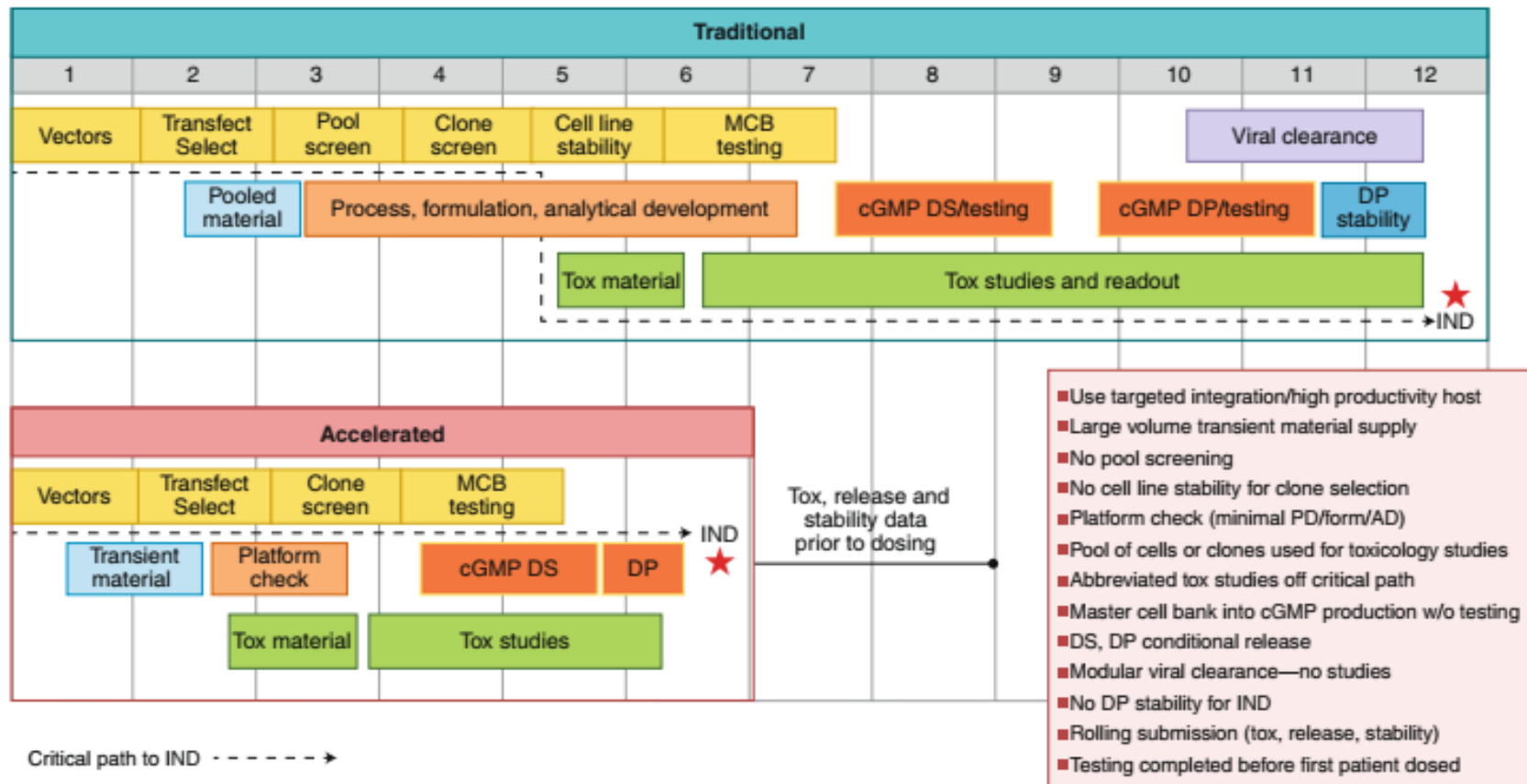
# Antigenic target for SARS-CoV-2 virus

- ▶ The primary antigenic epitope is the S protein
- ▶ S- protein facilitates target cell binding and fusion after binding the angiotensin-converting enzyme 2 (ACE2) receptor on the cell surface
- ▶ ACE2 is found on cells in the respiratory system, gastrointestinal tract and endothelium

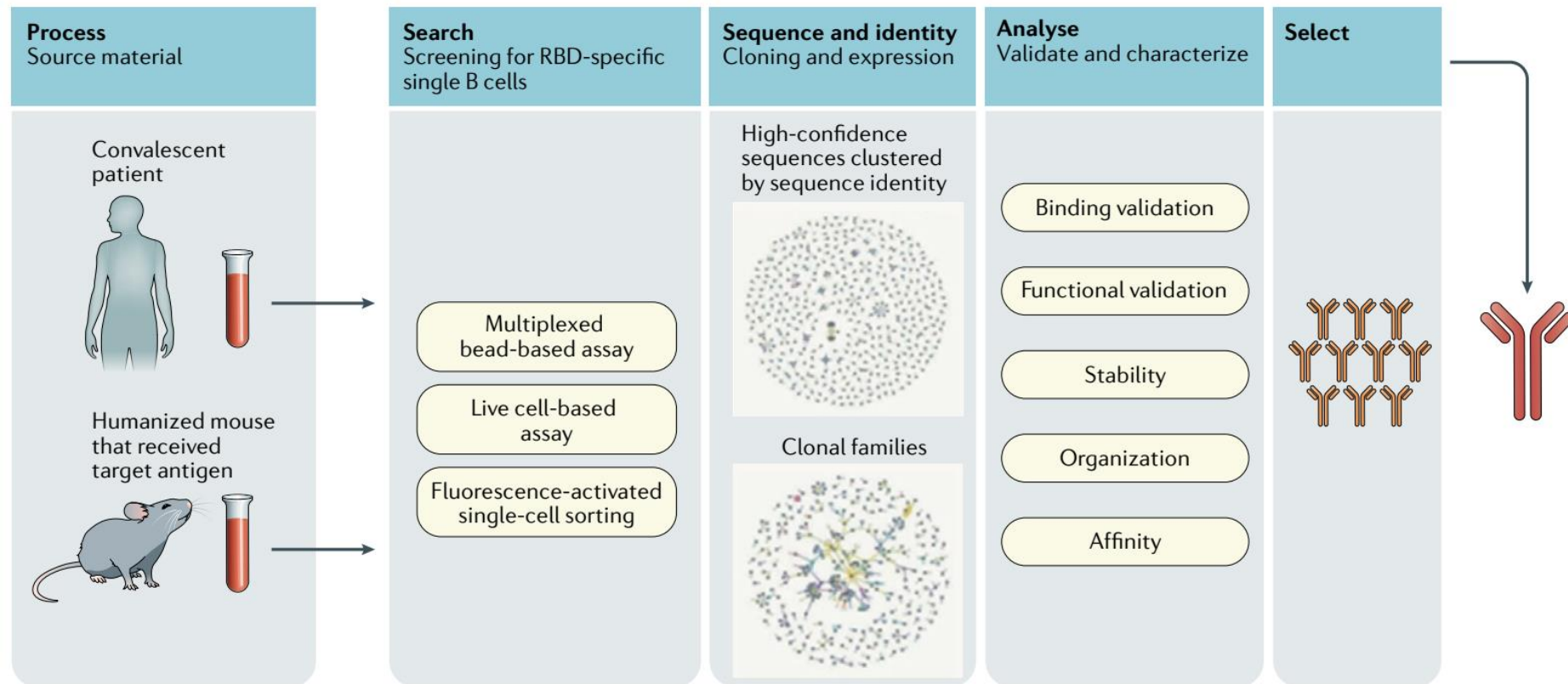
All antibodies designed to interact and engage S protein can neutralize the ability of the virus to bind and fuse with the target host cell



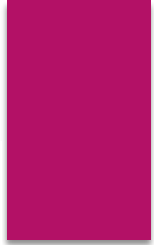
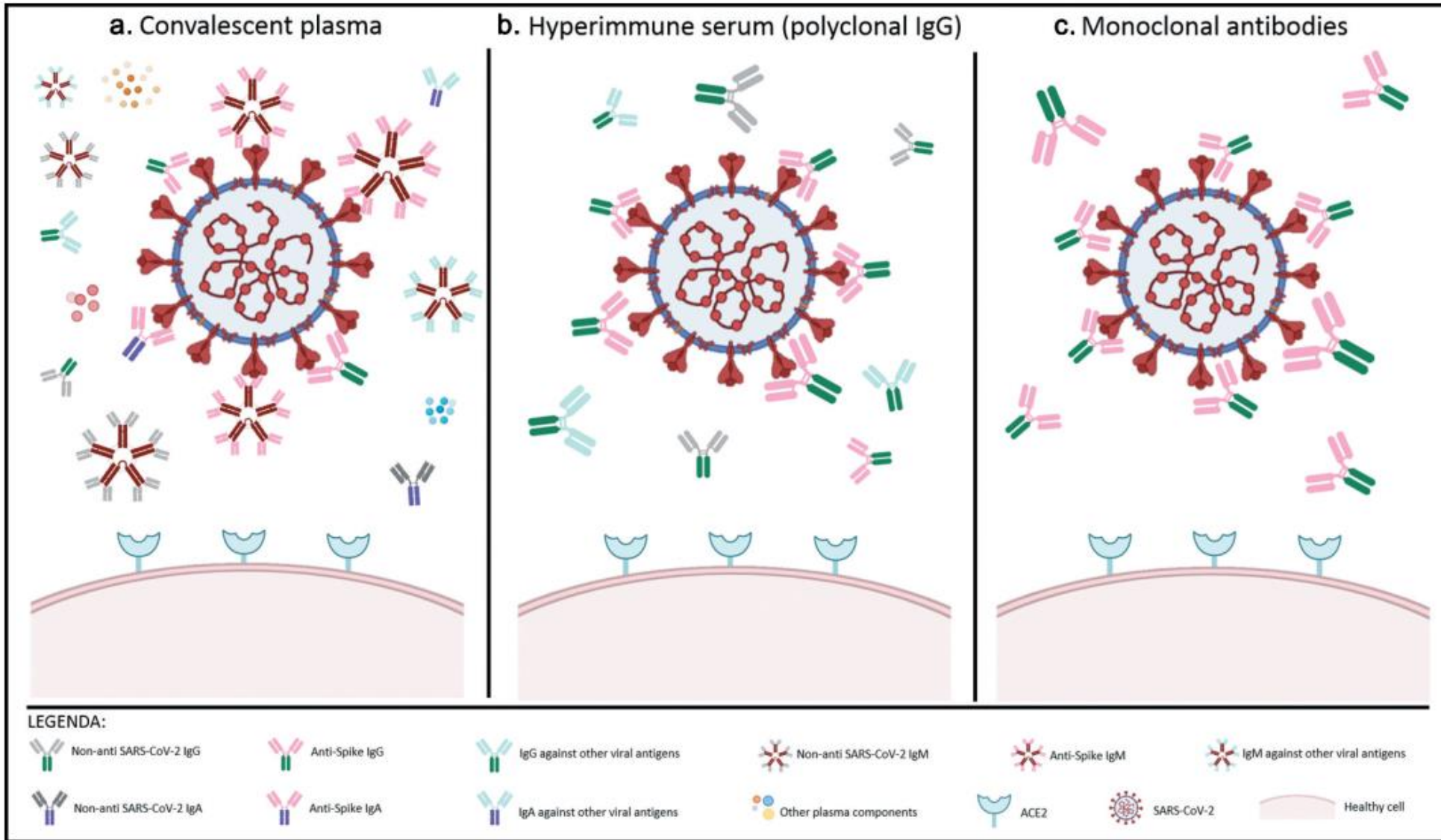
# Monoclonal antibodies design “usual” vs. “COVID” timeline

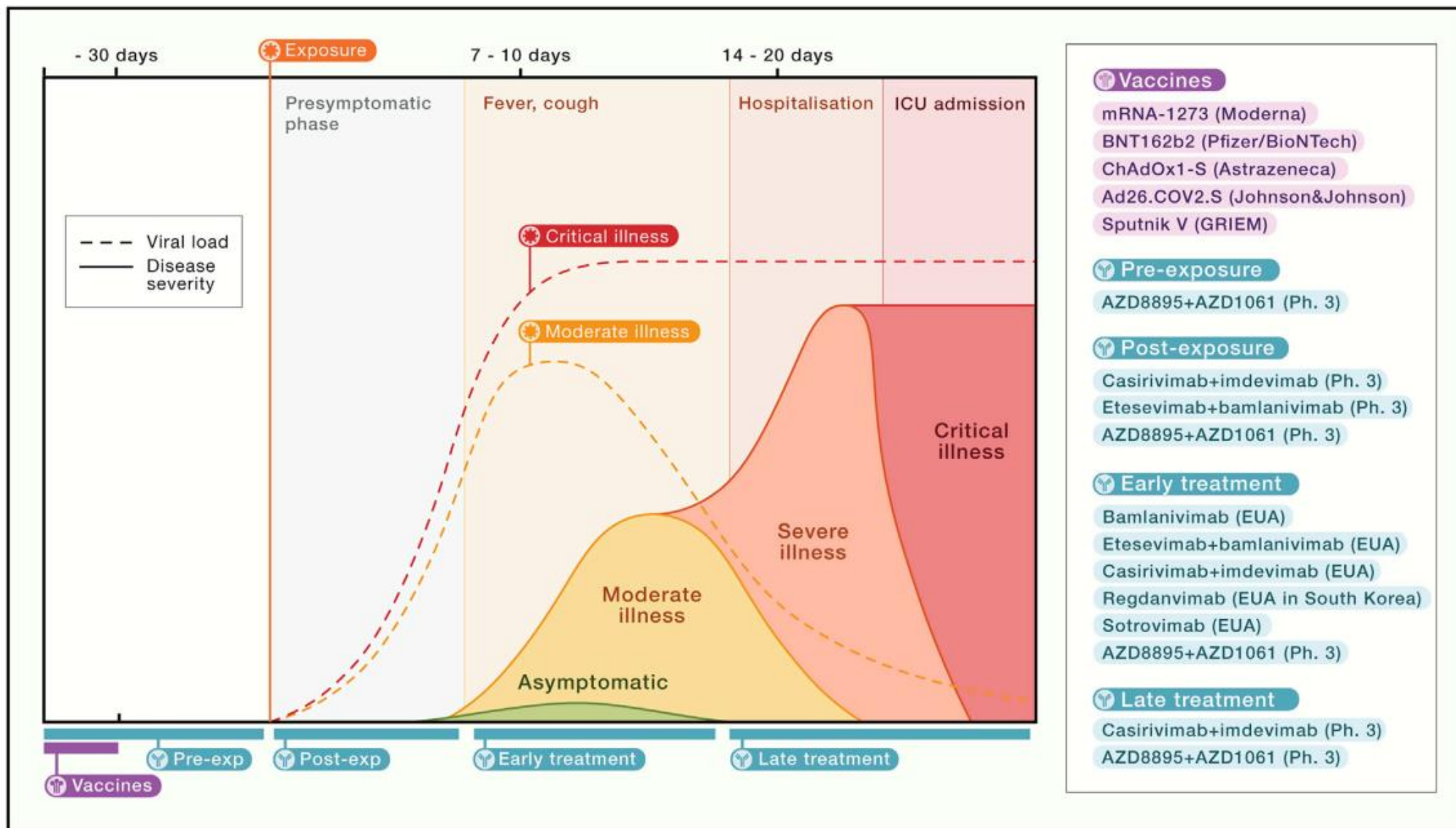


# Easiest way to obtain anti-SARS-CoV-2 mAbs

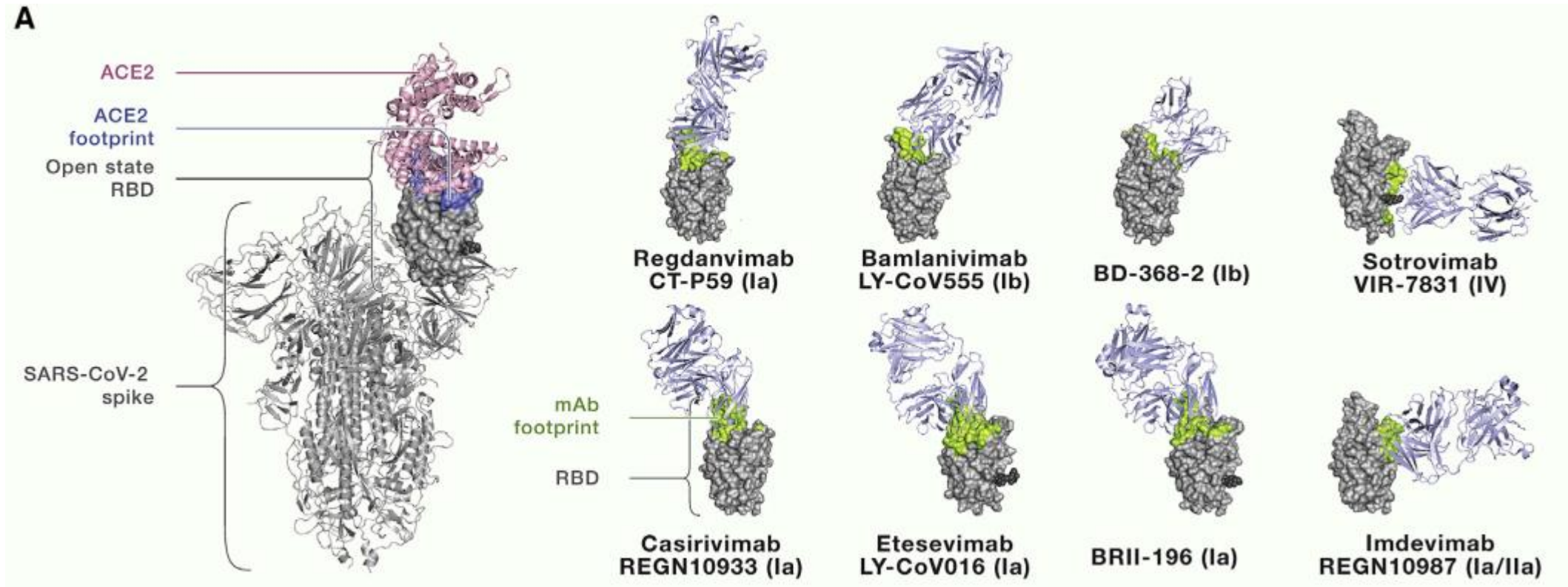








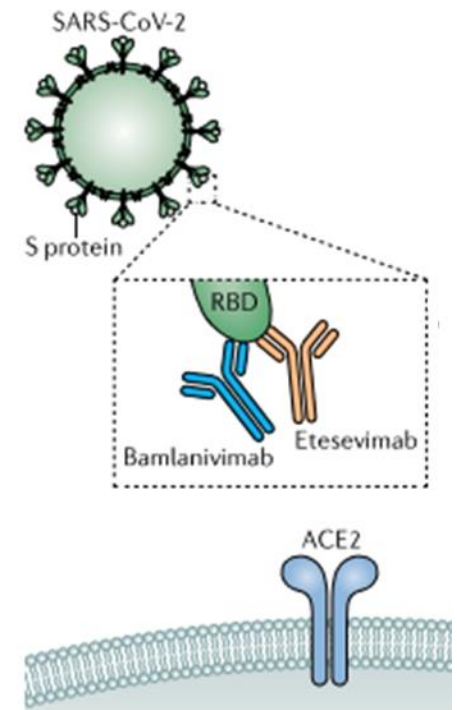
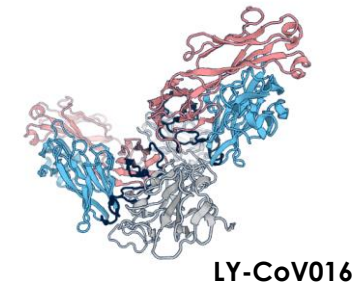
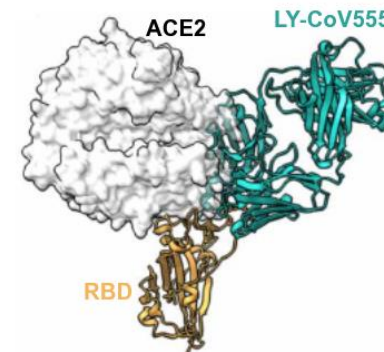
# Where/how is mAb connected to SARS-CoV 2





# BAMLANIVIMAB & ETESEVIMAB

- ▶ Bamlanivimab (LY-CoV555) was derived from the **convalescent plasma** of a patient who had COVID-19
  - binds the S protein's RBD, engaging its cognate epitope in both up and down conformations, which makes this antibody potentially useful as a **monotherapy**
  - single N-linked glycosylation site on each heavy chain
  - unmodified in the Fc region
- ▶ Etesevimab (LY-CoV016) presents amino acid substitutions in the Fc region (L234A, L235A) to **reduce effector function**



# BAMLANIVIMAB & ETESEVIMAB

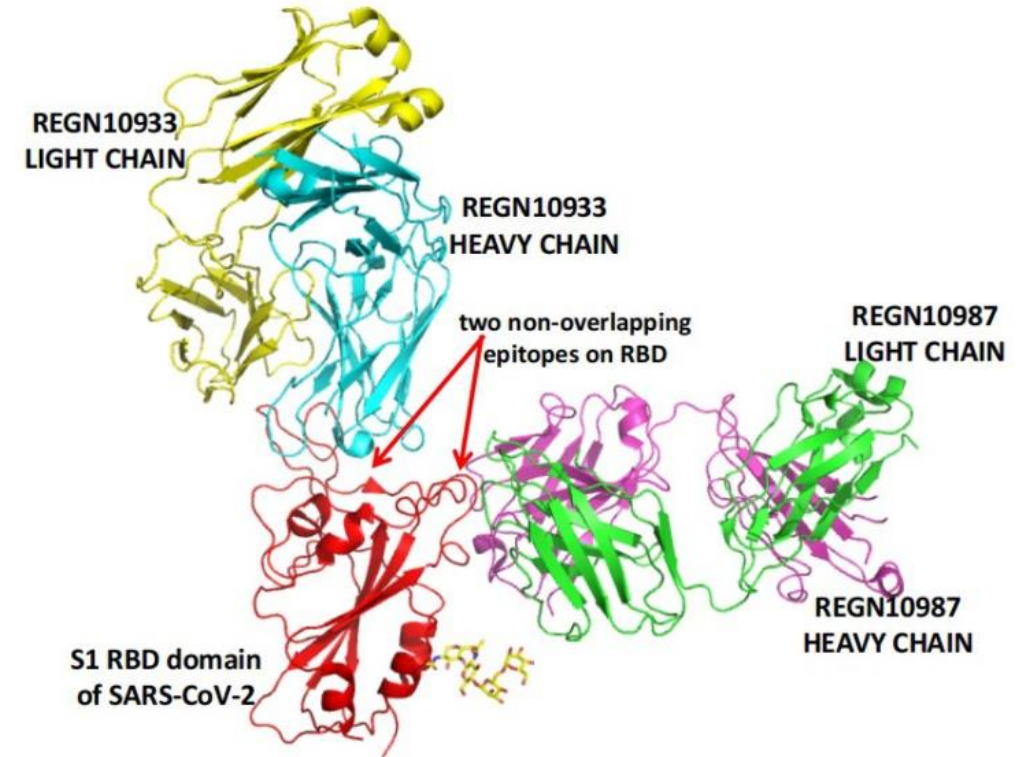
- ▶ Both exhibit neutralisation activity by blockage of the S protein interaction with angiotensin-converting enzyme 2 (ACE2) and prevent viral entry into human cells and viral replication targeting **overlapping, but different epitopes** of the RBD
- ▶ Produced in CHO cell lines
- ▶ Excipients/stabilizers include L-histidine buffer, sucrose, polysorbate 80, sodium chloride and WFI, and must be administered as a single intravenous infusion immediately after dilution with 0.9% sodium chloride





# REGN-COV2 (Casirivimab & Imdevimab )

- ▶ **Pooled** from > 200 neutralizing mAbs derived from convalescent plasma and infected humanized mice
- ▶ Casirivimab (REGN10933) and Imdevimab (REGN10987) are **IgG1** recombinant monoclonal antibodies targeting non-overlapping epitopes of the RBD of the S protein and have **unmodified Fc regions** and both contain single N-linked glycosylation site on each heavy chain
- ▶ REGN-COV2 exhibits neutralisation activity by blockage of the S protein interaction with angiotensin-converting enzyme 2 (ACE2), antibody-mediated cytotoxicity and cellular phagocytosis in virally infected cells was shown *in vitro*



# REGN-COV2 (Casirivimab & Imdevimab )

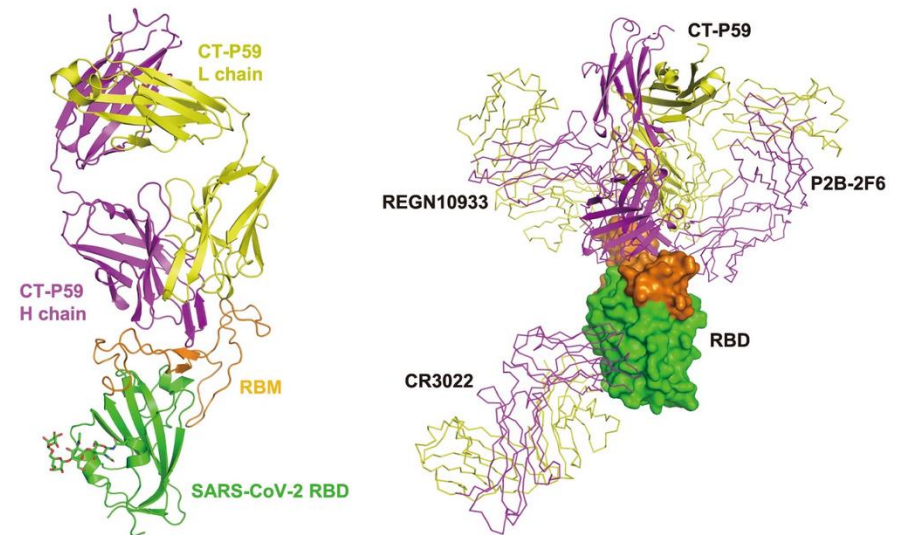
- ▶ Produced in CHO cell line
- ▶ Excipients/stabilizers: L-histidine and L-histidine monohydrochloride monohydrate (target pH 6.0), sucrose, polysorbate 80 and WFI, and is administered together as a single intravenous infusion immediately after dilution with 0.9% sodium chloride



# REGDANVIMAB

- ▶ Regdanvimab (CT-P59) recombinant human monoclonal IgG1 antibody targeting the RBD of the SARS-CoV-2 spike protein by blocking S protein interaction with ACE2
- ▶ contains a single N-linked glycosylation site on each heavy chain
- ▶ unable to mediate Fc-related activities

- binding orientation between CT-P59 and the RBD is different from other neutralizing mAbs
- CT-P59 can be a novel binder to the RBD
- CT-P59 has been proven to neutralize the D614G variant, one of the most infectious S protein's mutations



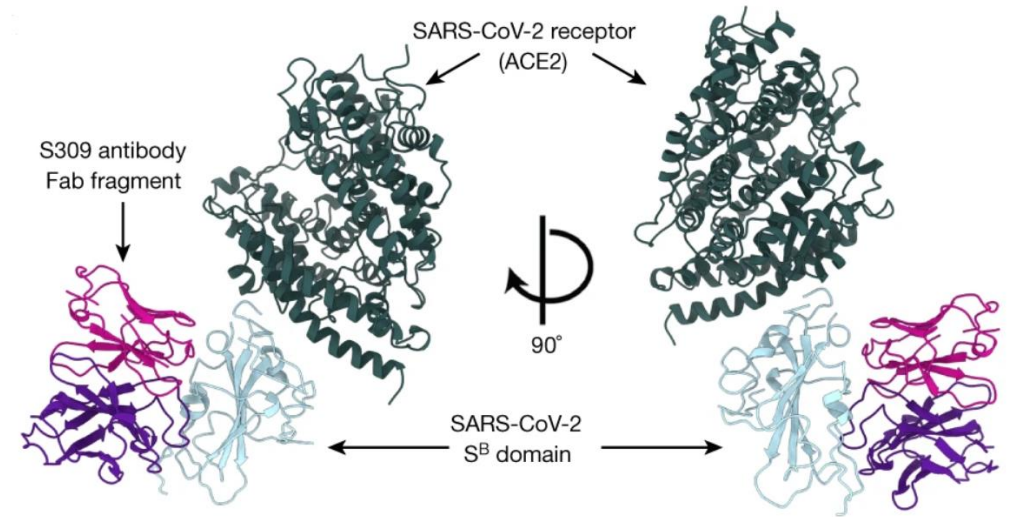
# REGDANVIMAB

- ▶ Produced in a CHO cell line
- ▶ Excipients/stabilizers - L-histidine and L-histidine hydrochloride monohydrate, L-arginine, polysorbate 80 and WFI



# SOTROVIMAB

- ▶ Sotrovimab (GSK4182136, VIR-7831) is an engineered human immunoglobulin monoclonal antibody of the IgG1κ subtype
- ▶ It is based on the identification and characterization of S309, an antibody identified from a **convalescent patient, who recovered from SARS-CoV in 2003**
- ▶ Sotrovimab binds to the SARS-CoV-2 RBD but can also engage in Fc-mediated receptor activities








































# SOTROVIMAB

- ▶ Produced in CHO cell line
- ▶ Excipients/stabilizers include L-histidine and L-histidine monohydrochloride buffer, L-methionine buffer, sucrose, polysorbate 80 and WFI



# Efficacy of neutralization vs. different strains of SARS-CoV-2

	B.1.1.7 (UK)	B.1.351 (South Africa)	P.1 (Brazil)	B.1.429 (California)	B.1.1.258 (Scotland)	B.1.525 (Nigeria)	B.1.526 (New York)	B.1.617.1 (India)
Casirivimab								
Imdevimab								
Bamlanivimab								
Etesevimab								
Sotrovimab								
Brii-196								
Brii-198								
AZD8895								
AZD1061								
Regdanvimab								
ADG-20								
BGB-DXP593								
ABBV-47D11								
ABBV-2B04								

 Neutralized  
( $<10$ -fold loss of neutralization)  
 Poorly or not-neutralized  
( $>10$ -fold loss of neutralization)  
 Predicted to be neutralized  
 Predicted to be weakly or to not be neutralized  
 Unknown  

\*Prediction of neutralization coverage is based on the presence of mutations in available epitope of each mAb

# FORMULATION of mAbs

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Antibody therapeutics are **large** (typically >150 kDa), **complex** in nature (most are glycosylated) and must be administered in **stoichiometric** rather than in catalytic quantities (nearly g/dose is required for many antibodies to be effective)

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An initial formulation effort will likely employ an **isotonic, non-hemolytic, slightly acidic** solution that can be administered by **intravenous** infusion or **subcutaneous** injection in preclinical studies

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Some stability information for the antibody drug can be obtained from initial *in silico* and *in vivo* testing during this preclinical phase that aids in the selection of a formulation that is used in clinical trials

# FORMULATION of mAbs

## Liquid sterile formulation

- ▶ Ready-to use
- ▶ Classical isotonic formulation with addition of stabilizers
- ▶ Less time to process

## Lyophilized (freeze-dried) formulation

- ▶ Needs to be reconstituted
- ▶ Increased stability due to low water content
- ▶ Transport advantages

# Stability of mAbs

Instability triggered by temperature change, pH change, water content, presence of salts and excipients, mechanical factors

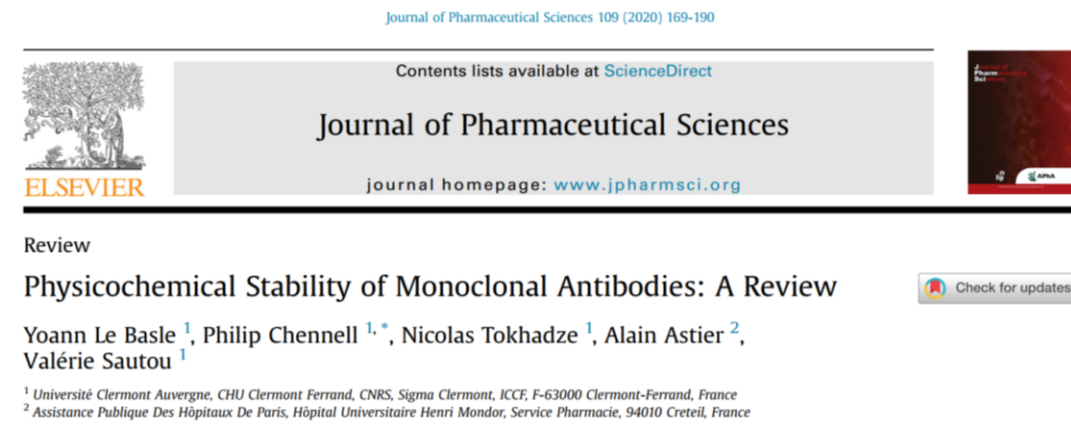
**mAbs instability may result in decrease of the biological activity and increase in immunogenicity**

## Physical instability

Aggregation  
Surface adsorption  
Denaturation

## Chemical instability

Deamidation  
Changes in disulphide bonds  
Isomerisation  
Oxydation  
Hydrolysis  
Fragmentation





# Challenges in formulation of protein pharmaceuticals

- ▶ Production process design
- ▶ Selection of containers (glass and plastic surfaces adsorb proteins and peptides)
- ▶ Careful selection of excipients
  - High risk of aggregation
  - Potential loss of activity
- ▶ Need of sterility for parenterals
- ▶ Stability and shelf-life

# What excipients are usually used?

- ▶ The basic amino acids histidine, lysine, and arginine stabilize all three domains of IgG
- ▶ Sugars and polyols are used as cryo-and lyoprotectants
- ▶ Surface active substances lower aggregation
- ▶ Different buffer mixtures are used in order to obtain desired pH/stability

Excipients	Frequency of Inclusion in MAb Formulations (%)		
	All	Lyophilized	Liquid
Polysorbate 80	57	45	62
Polysorbate 20	19	36	12
Poloxamer 188	3	0	4
Mannitol	8	9	8
Sorbitol	3	9	0
Sucrose	35	82	15
Trehalose	14	18	12
Dextrose	3	9	0
Dextran 40	3	9	0
NaCl	49	18	62
Arginine	8	0	12
Glycine	8	0	12
Methionine	3	0	4
Ascorbic acid	3	0	4
NaOAc	3	9	0
Phosphate	32	27	35
Citrate	11	9	12
Acetate	16	0	23
Tris	3	0	4
Succinate	3	9	0
Histidine	35	27	38

# Which excipients in anti- SARS-Cov-2 mAbs were used so far?

Anti-SARS-CoV2 mAbs	Excipients
Bamlanivimab and Etesevimab	L-histidine buffer, sucrose, polysorbate 80, sodium chloride and WFI
Casirivimab & Imdevimab	L-histidine and L-histidine monohydrochloride monohydrate (target pH 6.0), sucrose, polysorbate 80 and WFI
Regdanvimab	L-histidine and L-histidine hydrochloride monohydrate, L-arginine, polysorbate 80 and WFI
Sotrovimab	L-histidine and L-histidine monohydrochloride buffer, L-methionine buffer, sucrose, polysorbate 80 and WFI

# What future holds?

- ▶ The approved agents are all conventional full-length IgG
- ▶ More than 50 mAbs are still in clinical trials
- ▶ Apart from the antibody format, engineering has been accomplished in the natural isolates in different ways, e.g., the LALA mutation was introduced into the Fc portion of CB6 to lower the risk of Fc-mediated acute lung injury
- ▶ It is possible for IgM/IgA, IgY based therapeutics, bi- or tri-specific antibodies, singledomain- derived from phage display libraries, fusion proteins, etc. to soon progress in clinical trials
- ▶ Potential combining neutralizing mAb with the catalytic activity of Cas13, a fusion-protein targeting, and cutting the viral RNA (AntiBody And CAS fusion (ABACAS)), so that agents would exert both a prophylactic and anti-viral effect

# HVALA NA PAŽNJI

